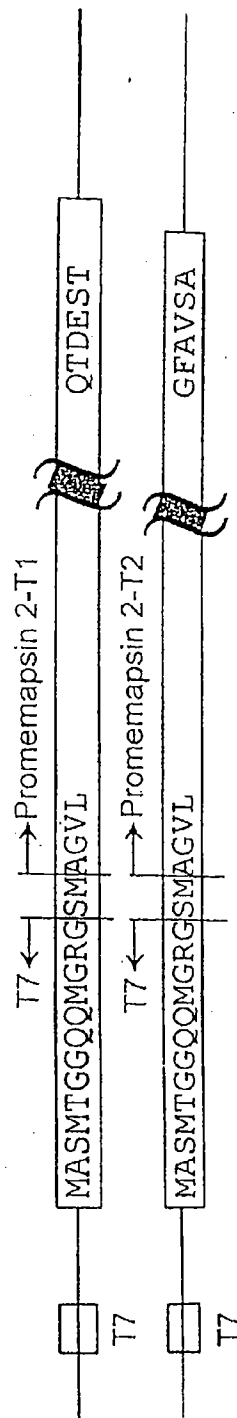


FIG. 1

1v MASMTGGQQM GRGSMAGVLP AHGTQHGIRL PLRSLGGAP LGLRLPRETD
 36p EEPPEPGRRG SFVEMVDNLR GKSGQGYVE MTVGSPQTL NILVDTGSSN
 38 FAVGAAPHF LHRYYQRQLS STYRDLRKG VVPYTQKWE GELGTDLVSI
 88 PHGPNVTVRA NIAAITESDK FFINGSNWEG ILGLAYAEIA RPDDSLPEFF
 138 DSLVKQTHVP NLFSLQLCGA GFPLNQSEVL ASVGSGMIIG GIDHSLYTGS
 188 LWYTPIRREW YYEVIIVRVE INGQDLKMDK KEYNYDKSIV DSGTTLRLP
 238 KKVFEAAVKS IKAASSTEKF PDGFWLGEQL VCWQAGTPW NIFPVISLYL
 288 MGEVTNQSER ITILPQQYLR PVEDVATSQD DCYKFAISQS STGTVMGAVI
 338 MEGFYVVFDR ARKRIGFAVS ACHVHDEFT AAVEGPFVTL DMEDCGYNIP
 388 QTDESTLMTI AYVMAAICAL FMLPLCLMVC QWRCLRLRQ QHDDFADDIS
 438 LLK*



T7 Protein (SEQ ID NO: 3)
 Memapsin 2T1 and 2T2 Construct (SEQ ID NOS: 32-34)